```
Sequence Listing could not be accepted due to errors.
See attached Validation Report.
If you need help call the Patent Electronic Business Center at (866)
217-9197 (toll free).
Reviewer: markspencer
Timestamp: [year=2009; month=11; day=18; hr=14; min=30; sec=11; ms=788;
Reviewer Comments:
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<213> Homo sapiens
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                                              205
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                                                               668
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   210
                       215
                                          220
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Codons must be made up of three nucleotides. The last codon in SEQ ID #
481 is only two nucleotides. Please make all necessary changes.
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               Artificial or Unknown found in <213> in SEQ ID (2)
W213
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Artificial or Unknown found in <213> in SEQ ID (3)

W213

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Artificial or Unknown found in <213> in SEQ ID (4)
W213
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W213
W213
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W213
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W213
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W213
                Artificial or Unknown found in <213> in SEQ ID (19)
W213
                Artificial or Unknown found in <213> in SEQ ID (20) This
W213
error has occured more than 20 times, will not be displayed
                Invalid sequence data feature in <221> in SEQ ID (636)
E257
                Invalid sequence data feature in <221> in SEQ ID (636)
E257
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E257
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E257
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E257
E257
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E257
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The warnings and errors shown above are ok and require no response.

Validated By CRFValidator v 1.0.3

Application No: 10045674 Version No: 4.0

Input Set:

Output Set:

Started: 2009-11-05 17:45:21.376

Finished: 2009-11-05 17:45:29.384

Elapsed: 0 hr(s) 0 min(s) 8 sec(s) 8 ms

Total Warnings: 459

Total Errors: 9

No. of SeqIDs Defined: 637

Actual SeqID Count: 637

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W	213	Artificial o	r	Unknown	found	in	<213>	in	SEQ	ID	(4)
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W	213	Artificial o	r	Unknown	found	in	<213>	in	SEQ	ID	(19)
W	213	Artificial o	r	Unknown	found	in	<213>	in	SEQ	ID	(20)

Input Set:

Output Set:

Started: 2009-11-05 17:45:21.376 **Finished:** 2009-11-05 17:45:29.384

Elapsed: 0 hr(s) 0 min(s) 8 sec(s) 8 ms

Total Warnings: 459
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No. of SeqIDs Defined: 637
Actual SeqID Count: 637

Error code		Error Description
		This error has occured more than 20 times, will not be displayed
E	320	Wrong Nucleic Acid Designator, gc in SEQID (481)
E	257	Invalid sequence data feature in <221> in SEQ ID (636)
E	257	Invalid sequence data feature in <221> in SEQ ID (636)
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SEQUENCE LISTING

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     COHEN, EDWARD H.
      NASTRI, HORACIO G.
      ROOKEY, KRISTIN L.
      HOET, RENE
      HOOGENBOOM, HENDRICUS R. J. M.
<120> NOVEL METHODS OF CONSTRUCTING LIBRARIES COMPRISING
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